

Fig. 1

HUMAN: 1 CAGACATCTGTGTCCCCCTCAAAAGTCATCCTGCCCCCGGGGAGGCTCCGTGCTGCTGACA
 CHIMPANZEE: |||||L|||TCTGTCTCCCCCAAAAGTCATCCTGCCCCCGGGGAGGCTCCGTGCTGCTGACA
 Q T S V S P P K V I L P R G G S V Q V T
 TGCAGCACCTCCTGTGACCAGCCCAAGTTGTTGGGCATAGAGACCCCGTTGCCCTAAAAAG
 |||||L|||TCTGTGACCAGCCCGACTTGTTCGGCCATAGAGACCCCGTTGCCCTAAAAAG
 TGCAGCACCTCCTGTGACCAGCCCGACTTGTTCGGCCATAGAGACCCCGTTGCCCTAAAAAG
 C S T S C D Q P D L L G I E T P L P K K
 HUMAN: 121 GAGTTGCTCCTGCTGGGAACAACCGGAAGGTGTATGAACCTGAGCAATGTGCAAGAAGAT
 CHIMPANZEE: |||||L|||TCTGCTTCTGGGTGGGAACAACCTGGAAGGTGTATGAACCTGAGCAATGTGCAAGAAGAT
 GAGTTGCTTCTGCTTCTGGGTGGGAACAACCTGGAAGGTGTATGAACCTGAGCAATGTGCAAGAAGAT
 E L L L G G N N W K V Y E L S N V Q E D
 AGCCAAACCAATGTGCTATTCAAACTGCCCTGATGGGCAGTCAACAGCTAAAAACCTTCCTC
 |||||L|||TCTGCTTCTGGGTGGGAACAACCTGCCCTGATGGGCAGTCAACAGCTAAAAACCTTCCTC
 AGCCAAACCAATGTGCTATTCAAACTGCCCTGATGGGCAGTCAACAGCTAAAAACCTTCCTC
 S Q P M C Y S N C P D G Q S T A K T F L
 HUMAN: 241 ACCGTGTACTGGACTCCAGAACGGGTGGAACTGGCACCCCTCCCTCTTGGCAGCCAGTG
 CHIMPANZEE: |||||L|||TCTGCTTCTGGGTGGGAACAACCTGGAAGGTGTATGAACCTGAGCAATGTGCAAGAAGAT
 ACCGTGTACTGGACTCCAGAACGGGTGGAACTGGCACCCCTCCCTCTTGGCAGCCAGTG
 T V Y W T P E R V E L A P L P S W Q P V
 GGCAAGAACCTTACCCCTACGCTGCCAGGTGGAGGGTGGGGCACCCCGGCCAACCTCACC
 |||||L|||TCTGCTTCTGGGTGGGAACAACCTGGAAGGTGTATGAACCTGAGCAATGTGCAAGAAGAT
 GGCAAGAACCTTACCCCTACGCTGCCAGGTGGAGGGTGGGGCACCCCGGCCAACCTCACC
 G K D L T L R C Q V E G A P R A N L T
 HUMAN: 361 GTGGTGCTGCTCCGTGGGAGAAAGGAGCTGAAACGGGAGCCAGCTGTGGGGGAGCCCGCT
 CHIMPANZEE: |||||L|||TCTGCTTCTGGGTGGGAACAACCTGGAAGGTGTATGAACCTGAGCAATGTGCAAGAAGAT
 GTGGTGCTGCTCCGTGGGAGAAAGGAGCTGAAACGGGAGCCAGCTGTGGGGGAGCCCGCT
 V V L L R G E K E L K R E P A V G E P A
 GAGGTCACGACCAACCGGTGCTGGTGAGGAGAGATCACCATGGAGCCCAATTCTCGTGCCGC
 |||||L|||TCTGCTTCTGGGTGGGAACAACCTGGAAGGTGTATGAACCTGAGCAATGTGCAAGAAGAT
 GAGGTCACGACCAACCGGTGCTGGTGAGGAGAGATCACCATGGAGCCCAATTCTCGTGCCGC
 E V T T T V L V E R D H H G A N F S C R

FIG. 2

HUMAN: 481 ACTGAACTGGACCTGCGGCCCCCAAGGGCTGGAGCTGTTTGAGAACACCTCGGCCCTTAC
 CHIMPANZEE: T E L D L R P Q G L Q L F E N T S A P H
 CAGCTCCAGACCTTTGTCCCTGCCAGCGACTCCCCCACAACTTGTTCAGCCCCCGGGTCCCTA
 CAGCTCCAAACCTTTGTCCCTGCCAGCGACTCCCCCACAACTTGTTCAGCCCCCGGGTCCCTA
 Q L Q T F V L P A T P Q L V S P R V L
 HUMAN: 601 GAGGTGGACACGCAGGGGACCGTGGTCTGTTCCTGGACGGGCTGTTCCTCAGTCTCGGAG
 CHIMPANZEE: GAGGTGGACACGCAGGGGACCGTGGTCTGTTCCTGGAGTGGGCTGTTCCTCAGTCTTGGAG
 E V D T Q G T V C S L D G L F P V L E
 GCCCAGGTCCACCTGGCACCTGGGGGACCCAGAGGTTGAAACCCACAGTCACCTATGGCAAC
 GCCCAGGTCCACCTGGCACCTGGGGGACCCAGAGGTTGAAACCCACAGTCACCTATGGCAAT
 A Q V H L A L G D Q R L N P T V T Y G N
 HUMAN: 721 GACTCCTTCTCGGCCCAAGGCCCTCAGTCAGTGTGACCCGACAGGACGAGGACCCAGCGG
 CHIMPANZEE: GACTCCTTCTCGGCCCAAGGCCCTCAGTCAGTGTGACCCGACAGGACGAGGACCCAGCGG
 D S F S A K A S V S V T A E D E G T Q R
 CTGACGTGTGCAGTAATACTGGGGAACCCAGAGCCAGGACACACTGCAGACAGTGACCATC
 CTGACGTGTGCAGTAATACTGGGGAACCCAGAGCCGAGACACACTGCAGACAGTGACCATC
 L T C A V I L G N Q S R E T L Q T V T I
 HUMAN: 841 TACAGCTTTCGGGCGCCCAACGTGATTCTGACGAAGCCAGAGGTCTCAGAAGGACCCGAG
 CHIMPANZEE: TACAGCTTTCGGGCGCCCAACGTGATTCTGACGAAGCCAGAGGTCTCAGAAGGACCCGAG
 Y S F P A P N V I L T K P E V S E G T E
 GTGACAGTGAAGTGTGAGGCCCAACCCCTAGAGCCCAAGGTGACCGCTGAATGGGGTTCAGCCC
 GTGACAGTGAAGTGTGAGGCCCAACCCCTAGAGCCCAAGGTGACCGCTGAATGGGGTTCAGCCC
 V T V K C E A H P R A K V T L N G V P A

FIG. 2 (CONT.)

HUMAN:	961	CAGCCACTGGGCCCCGAGGGCCCCAGCTCCTGCTGAAGGCCACCCAGAGGACAAACGGGGCC
CHIMPANZEE:		Q P V G P R V Q L L K A T P E D N G R
		AGCTTCTCCTGCTCTGCAACCCCTGGAGGTGGCCCGCCAGCTTATACACAAGAACCCAGACC
		AGCTTCTCCTGCTCTGCAACCCCTGGAGGTGGCCCGCCAGCTTATACACAAGAACCCAGACC
		S F S C S A T L E V A G Q L I H K N Q T
HUMAN:	1081	CGGGAGCTTCGTCTCCTGTATGGCCCCCGACTGGACGAGAGGGATTGTCCGGGAAACTGG
CHIMPANZEE:		R E L R V L Y G P R L D E R D C P G N W
		ACGTGGCCAGAAATTCCCGAGCAGACTCCCAATGTGCCAGGCTTGGGGGAACCCATTGCCCC
		ACGTGGCCAGAAATTCCCGAGCAGACTCCCAATGTGCCAGGCTTGGGGGAACCCATTGCCCC
		T W P E N S Q Q T P M C Q A S G N P L P
HUMAN:	1201	GAGCTCAAGTGCTCTAAAGGATGGCACTTTCCCACTGCCCACTCGGGGAATCAGTGACTGTC
CHIMPANZEE:		E L K C L K D G T F P L P V G E S V T V
		ACTCGAGATCTTGAGGGCACCTACCTCTGTCTGGGGCCAGGAGCACTCAAGGGAGGTCACC
		ACTCGAGATCTTGAGGGCACCTACCTCTGTCTGGGGCCAGGAGCACTCAAGGGAGGTCACC
		T R D L E G T Y L C R A R S T Q G E V T
HUMAN:	1321	CGCGAGGTGACCGTGAAATGTGCTCTCCCCCGGTATGAGATTGTTCATCATCTGTGCTA
CHIMPANZEE:		R K V T V N V L S P R Y E I V I T V V
		GCAGCCGCAGTCATAATGGGCACCTGCAGGCCCTCAGCACCTACCTCTATAACCCGCCAGGG
		GCAGCCGCAGTCATAATGGGCACCTGCAGGCCCTCAGCACCTACCTCTATAACCCGCCAGGG
		A A V I M G T A G L S T Y L Y N R Q R

FIG. 2 (CONT.)

HUMAN:	1441	AAGATCAAGAAATACAGACTACAAACAGGCCCAAAAGGGA	CCCCCATGAAACCGAACACA
CHIMPANZEE:		AAGATCAGGAAATACAGACTACAAACAGGCTCAAAAGGGA	CCCCCATGAAACCGAACACA
		K I R K Y R L Q Q A Q K G T P M K P N T	
		CAAGCCACGCCCTCCCTGA	
		CAAGCCACGCCCTCCCTGA	
		Q A T P P ^ ^ ^	

FIG. 2 (CONT.)

1515 ICAM
 CAG ACA TCT GTG TCC CCC CCA AAA GTC ATC CTG CCC CGG GGA GGC TCC GTG CTG GTG ACA
 TGC AGC ACC TCC TGT GAC CAG CCC ACC TTT TTT GGC ATA GAG ACC CCG TTG CCT AAA AAG
 GAG TTG CTC CTG CTT GGG AAC AAC CAG AAG GTG TAT GAA CTG AGC AAT GTG CAA GAA GAT
 AGC CAA CCA ATG TGT TAT TCA AAC TGC CCT GAT GGG CAG TCA ACA GCT AAA ACC TTC CTC
 ACC GTG TAC TGG ACT CCA GAA CGG GTG GAG GGT GGC GCA CCC TCT TGG CAG CCA GTG
 GGC AAG GAC CTT ACC CTA CGC TGC CAG GTG GAG GGT GGC GCA CCC GCT GTG GGC ACC GTC
 GTG GTG CTC CGT GGG GAG GAG CAG CAG AAA GAT CAC CAT GGA GCC AAT TTC TTG TGC CGC
 GAG GTC ACG ACC ACG GTG CCG GTG CAA GGG CTG AAG CTG TTT GAG AAC ACC TCG GCC CCC TAC
 ACT GAA CTG GAC CTG CGG CCC CAA GGG CTG AAG CTG CCA CAA CTT GTC AGC CCT GCG GAG
 CAG CTC CAA ACC TTT GTC CTG CCA GCG ACT GTG GGC GAC CAG AGG TTG AAC CCC ACA GTC ACC TAT GGC AAC
 GAG GTG GAC ACG CAG GGG ACT GTG GGC GAC CAG AGG TTG AAC CCC ACA GAG GAC AAC GGC CGC
 GCC CAG GTC CAC CAG GGC ACT GTG GGC GAC CAG AGG TTG AAC CCC ACA GAG GAC AAC GGC CGC
 GAC TCC TTC TCA GCC AAG GCC GTC GTC ACC GCA GAG GAC GAG ACA CTG CAG ACA GTG ACC ATC
 CTG ACG TGT GCA GTA ATA CTG GGG ACC CAG AGC CAG GAG ACA CTG CAG ACA GAG GGC ACC GAG
 TAC AGC TTT CCG GCA CCC AAC GTG ATT CTG ACG AAG CCA GAG GTC TCA GAA GGC ACC GGC
 GTG ACA GTG AAG TGT GAG GCC CAC CCT AGA GCC AAG GTG ACA CTG AAT GGC GGT CCA GCC
 CAG CCA CCG GGC CCG AGG ACC CAG TTC CTG CTG AAG GCC ACC CCA GAG AAC GGC CGC
 AGC TTC TCC TGT GCA ACC CTG GAG GTG GGC CCG CAG GTG GAT GAG AGG GAT TGT CCG GAA AAC TGG
 CGG GAG CTT CGT GTC TCC CAG CAG ACT CCA ATG TGC CAG GCT TGG GGC AAC CCA TTG CCC
 ACG TGG CCA GAA AAT TCC CAG CAG ACT TTC CCA CTG CCC GTC GGC GAA TCA GTG ACT GTC
 GAG CTC AAG TGT CTA AAG GAT GGC ACT TAC CTC TGT CGG GCC AGG AGC ACT CAA GGC GAG GTC ACC
 ACT CGA GAT CTT GAG GGC ACC TAC CTC TGT CGG GCC AGG AGC ACT CAA GGC GAG GTC ACC
 CGC GAG GTG ACC GTG AAT GTG CTC TCC CCC CGG TAT GAG TTT GTC ATC ATC GCT GTG GTA
 GCA GCC GCA ATA ATG GGC ACT GCA GGC CTC AGC ACG TAC CTC TAT AAC CGC CAG CGG
 AAG ATC AGG AAA TAC AGA CTA CAA CAG GCT CAA AAA GGC ACC CCC ATG AAA CCG AAC ACA
 CAA GCC ACG CCT CCC

GORILLA

(SEQ ID NO: 4)

Fig. 3

Fig. 4

Human J03132	QTSVSPSKVI	LPRGGSVLVT	CSTSCDQPKL	LGIETPLPKK	ELLPLGNRRK
Human X06990
Human X59286-8
Human #4
Human #7
Human #8
Human M24283
Human U86814M.
Chimp M86848P...Q..D.G...W.
Chimp #1P...Q..D.G...W.
Gorilla #1P...T.L...Q.
Gorilla #2P...T.L...Q.
Orang	H....SAN.FNT.PG...W.
Human J03132	VYELSNVQED	SQPMCYSNCP	DGQSTAKTFL	TVYWTPERVE	LAPLPSWQPV
Human X06990
Human X59286-8
Human #4
Human #7
Human #8
Human M24283
Human U86814
Chimp M86848
Chimp #1
Gorilla #1
Gorilla #2
Orang	M.....A...
Human J03132	GKNLTLRCQV	EGGAPRANLT	VVLLRGEKEL	KREPAVGPEA	EVTTLVLVRR
Human X06990
Human X59286-8
Human #4
Human #7
Human #8

(SEQ ID NO:6)

Fig. 5A

Human M24283
Human U86814	????????	????????	????????	????????	????????	????????	????????
Chimp M86848	.D.....E.
Chimp #1	.D.....E.
Gorilla #1	.D.....IE.P.EKP.EK
Gorilla #2	.D.....IE.P.EKP.EK
OrangE.	S.Q.....A...A.KA...A.K
Human J03132	DHHGANFSCR	TELDLRPQGL	ELFENTSAPY	QLQTFVLPTAT	PPQLVSPRVL		
Human X06990
Human X59286-8
Human #4
Human #7
Human #8
Human M24283
Human U86814	????????	????????	????????	????????	????????	????????	????????
Chimp M86848	Q.....H
Chimp #1	Q.....H
Gorilla #1L..	K.....
Gorilla #2L..	K.....
Orang	.D.....H
Human J03132	EVDTQGTVC	SLDGLFPVSE	AQVHLALGDQ	RLNPTVTYGN	DSFSAKASVS		
Human X06990
Human X59286-8
Human #4
Human #7
Human #8
Human M24283
Human U86814	????????	????????	????????	????????	????????	????????	????????
Chimp M86848L.
Chimp #1L.
Gorilla #1

Fig. 5B

Human X59286-8
Human #4
Human #7
Human #8
Human M24283
Human U86814	??????????	??????????	??????????	??????????	??????????	??????????	??????????	??????????	??????????
Chimp M86848S.....
Chimp #1S.....
Gorilla #1
Gorilla #2
Orang
Human J03132	ELKCLKDGTG	PLPIGESVT	TRDLEGTYLC	RARSTQGEVT	REVTVNVLS				
Human X06990
Human X59286-8
Human #4
Human #7
Human #8
Human M24283K.....
Human U86814	??????????	??????????	??????????	??????????	??????????	??????????	??????????	??????????	??????????
Chimp M86848V.....K.....
Chimp #1V.....K.....
Gorilla #1V.....
Gorilla #2V.....
Orang
Human J03132	RYEIVITVV	AAAVIMGTAG	LSTLYNQR	KIKYRLQA	QKGTMPKNT				
Human X06990
Human X59286-8
Human #4
Human #7
Human #8
Human M24283
Human U86814	??????????	??????????	??????????	??????????	??????????	??????????	??????????	??????????	??????????
Chimp M86848
Chimp #1

Fig. 5D

Gorilla #1	...F...A..R.....
Gorilla #2	...F...A..R.....
OrangA.L....R.....
Human J03132	QATPP				
Human X06990				
Human X59286-8				
Human #4				
Human #7				
Human #8				
Human M24283				
Human U86814	?????				
Chimp M86848				
Chimp #1				
Gorilla #1				
Gorilla #2				
Orang	.T...				

Fig. 5E

Human M32331	SDEKVFEVHV	RPKKLAVEPK	GSLEVNCSTT	CNQPEVGGLE	TSLDKILLDE
Human #4
Human #8
Human X15606N.....
Chimp #1K.....
Chimp #2K.....
Gorilla #2	A.....
Human M32331	QAQWKHYLVS	NISHDTVLQC	HFTCSGKQES	MNSNVSVYQP	PRQVILTLPQ
Human #4
Human #8
Human X15606
Chimp #1
Chimp #2
Gorilla #2
Human M32331	TLVAVGKSFT	IECRVPTVEP	LDLTLFLFR	GNETLHYETF	GKAAPAPQEA
Human #4
Human #8
Human X15606
Chimp #1
Chimp #2
Gorilla #2NQ..L...
Human M32331	TATFNSTADR	EDGHRNFSCL	AVLDLMSRGG	NIFHKHSAPK	MLEIYEPVSD
Human #4
Human #8
Human X15606
Chimp #1	.V.....	D.....
Chimp #2	.V.....	D.....
Gorilla #2I....	...QE.....

(SEQ ID NO:7)

Fig. 6A

Human M32331	SQMVIIVTVV	SVLLSLFVTS	VLLCFIFGQH	LRQQRMGTYG	VRAAWRRLPQ
Human #4
Human #8
Human X15606
Chimp #1
Chimp #2
Gorilla #2
Human M32331	AFRP				
Human #4				
Human #8				
Human X15606				
Chimp #1				
Chimp #2				
Gorilla #2				

Fig. 6B

Human X69819	QEFLLRVEPQ	NPVLSAGGSL	FVNCSTDCPS	SEKIALETSL	SKELVASGMG
Human #4
Human #5
Human #7
Human S50015	F.....
Chimp #3
Chimp #4
Chimp #5
Gorilla #1
Gorilla #2
OrangP....	L.....	.K.....DN...
Human X69819	WAAFNL SNVT	GNSRILCSVY	CNGSQITGSS	NITVYGLPER	VELAPLPPWQ
Human #4
Human #5
Human #7
Human S50015
Chimp #3R....
Chimp #4R....
Chimp #5R....
Gorilla #1R....
Gorilla #2R....
OrangY.....I...R....L..
Human X69819	PVGQNFTLRC	QVEGGSPRTS	LTVVLLRWEE	ELSRQPAVEE	PAEVTATVLA
Human #4
Human #5
Human #7
Human S50015
Chimp #3	Q.....
Chimp #4	Q.....
Chimp #5	R.....
Gorilla #1P...
Gorilla #2P...

(SEQ ID NO:8)

Fig. 7A

Human X69819	SRDDHGAPFS	CRTELDMQPQ	GLGLFVNTSA	PRQLRTFVLP	VTPPRLVAPR
Human #4
Human #5
Human #7
Human S50015
Chimp #3
Chimp #4
Chimp #5
Gorilla #1	..G.....	M.....
Gorilla #2	..G.....	M...S....
Orang	..GH...H..
Human X69819	FLEVETSWPV	DCTLDGLFPA	SEAQVYLALG	DQMLNATVMN	HGDTLTATAT
Human #4
Human #5
Human #7
Human S50015
Chimp #3
Chimp #4
Chimp #5
Gorilla #1
Gorilla #2
Orang	...A.....V.
Human X69819	ATARADQEGA	REIVCNVTLG	GERREARENL	TVFSFLGPIV	NLSEPTAHEG
Human #4
Human #5
Human #7
Human S50015
Chimp #3T.P..

Fig. 7B

Chimp #4T.P..
Chimp #5T.P..
Gorilla #1	...L.....I.....P..
Gorilla #2	...L.....I.....P..
Orang	.M.....	Q.....LS.P..
Human X69819	STVTVSCMAG	ARVQVTLDGV	PAAAPGQPAQ	LQLNATESDD	GRSFFCSATL
Human #4
Human #5
Human #7
Human S50015
Chimp #3	R.....
Chimp #4	R.....
Chimp #5	R.....
Gorilla #1
Gorilla #2
Orang
Human X69819	EVDGEFLHRN	SSVQLRVLYG	PKIDRATCPQ	HLKWKDKTRH	VLQCQARGNP
Human #4
Human #5
Human #7
Human S50015
Chimp #3T.
Chimp #4T.
Chimp #5T.
Gorilla #1T.
Gorilla #2T.
OrangF...
Human X69819	YPELRCLKEG	SSREVPVGIP	FFVNVTHNGT	YQCQASSSRG	KYTLVVVMDI
Human #4
Human #5
Human #7

Fig. 7C

Human S50015
Chimp #3
Chimp #4
Chimp #5
Gorilla #1
Gorilla #2
Orang	H.....	R.....
Human X69819	EAGSSHFVPV	FVAVLLTLGV	VTIVLALMYV	FREHQRSQSY	HVREESTYLP
Human #4
Human #5T.....
Human #7
Human S50015
Chimp #3K.....
Chimp #4K.....
Chimp #5K.....
Gorilla #1K.....
Gorilla #2K.....
Orang	...N...L.	.L...V...	.V.V.....K...R.	...Q...S..
Human X69819	LTSMQPTEAM	GEEPSRAE			
Human #4			
Human #5			
Human #7			
Human S50015			
Chimp #3Q..			
Chimp #4Q..			
Chimp #5			
Gorilla #1			
Gorilla #2			
OrangT..			

Fig. 7D

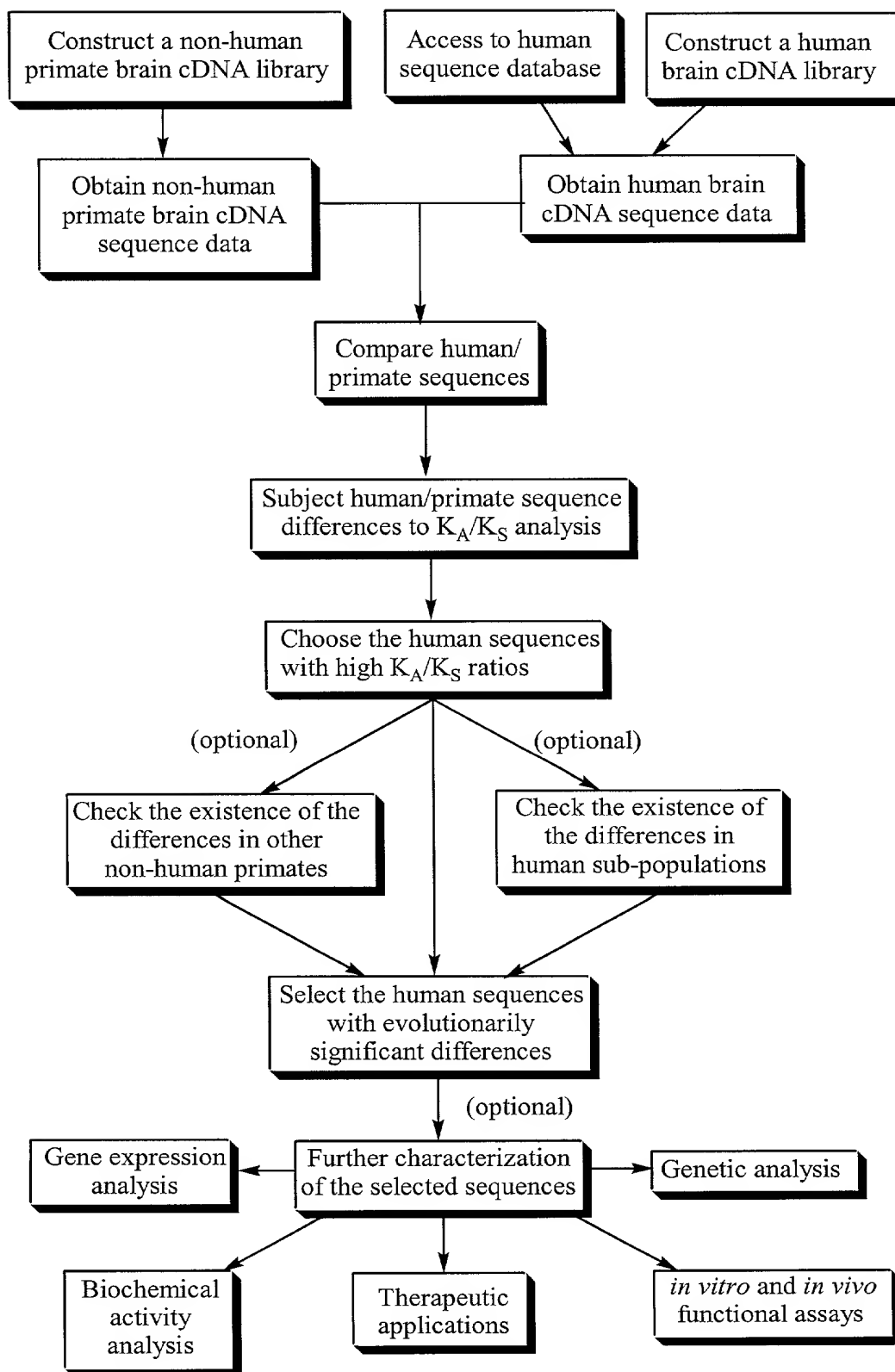


Fig. 8

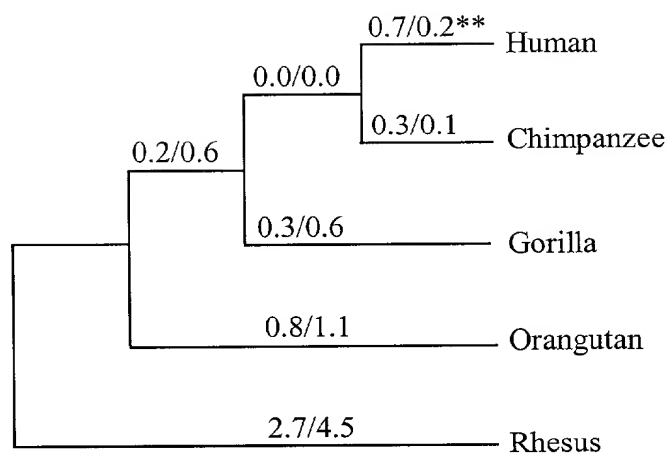


Fig. 9

094222.03280.1

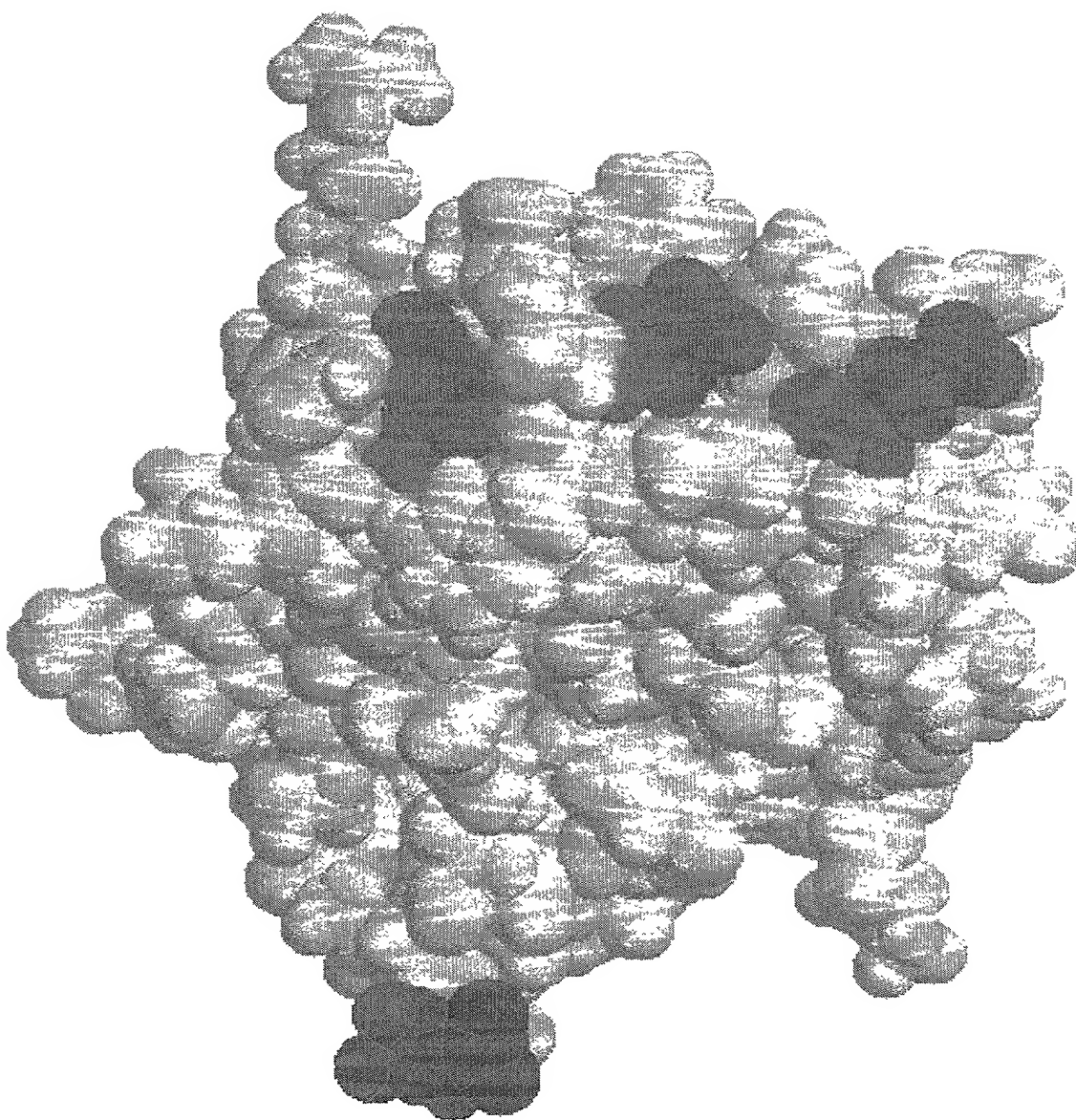


Fig. 10

Human

ATGAGTGACTCCAAGGAACCAAGACTGCAGCAGCTGGGCTCTCTGGAGGAGGAACA
GCTGAGAGGCCCTTGGAATCCGACAGACTCGAGGATACAAAGAGCTTAGCAGGGTGTC
TTGGCCATGGTCCCTGGTGCTGCAACTCCTCTCTCACGCTCTTGCTGGCTGGCTCCT
TGTCCAAAGTGTCCAAAGGTCCCCAGCTCCATAAGTCAGGAACAATCCAGGCAAGACG
CGATCTACCAGAACCTGACCCAGCTTAAAGCTGCAGTGGGTGAGCTCTCAGAGAAA
TCCAAGCTGCAGGAGATCTACCAGGAGCTGACCCAGCTGAAGCTGCAGTGGGTGA
GCTTCCAGAGAAATCTAAGCTGCAGGAGATCTACCAGGAGCTGACCCGGCTGAAGG
CTGCAGTGGGTGAGCTTCCAGAGAAATCTAAGCTGCAGGAGATCTACCAGGAGCTG
ACCTGGCTGAAGGCTGCAGTGGGTGAGCTTCCAGAGAAATCTAAGATGCAGGAGAT
CTACCAGGAGCTGACTCGGCTGAAGGCTGCAGTGGGTGAGCTTCCAGAGAAATCTA
AGCAGCAGGAGATCTACCAGGAGCTGACCCGGCTGAAGGCTGCAGTGGGTGAGCTT
CCAGAGAAATCTAAGCAGCAGGAGATCTACCAGGAGCTGACCCGGCTGAAGGCTGC
AGTGGGTGAGCTTCCAGAGAAATCTAAGCAGCAGGAGATCTACCAGGAGCTGACC
CAGCTGAAGGCTGCAGTGGAAACGCTGTGCCACCCCTGTCCCTGGGAATGGACATT
CTTCCAAAGGAAACTGTACTTCACTGTCTAACTCCAGCGGAACCTGGCACGACTCCAT
CACCGCTGCAAGAAAGTGGGGGCCAGCTCGTCTAATCAAAAGTGTGAGGAGC
AGAACTTCTACAGCTGCAGTCTTCCAGAAAGTAAACCGCTTCACTGGGACTTT
CAGATCTAAATCAGGAAGGCACGTGGCAATGGGTGGACGGCTCACCTCTGTGGCCC
AGCTTCAAGCAGTATTGGAAACAGAGGAGAGCCCAACAACGTTGGGAGGAAGACTG
CGCGGAATTTAGTGGCAATGGCTGGAACGACGACAAATGTAACTTGGCCAAATCTG
GATCTGCAAAAAGTCCGAGCTCTGCTCCAGGATGAAGAACAGTTTCTTCTCC
AGCCCTGCCACCCCAACCCCTCTCTCG (SEQ. ID. NO. 9)

Fig. 11

Chimpanzee

ATGAGTGACTCAAAGGAACCAAGACTGCAGCAGCTGGGCCCTCCTGGAGGAGGAACA
GCTGAGAGGCCCTTGGAATCCGACAGACTCGAGGCTACAAGAGCTTAGCAGGGTGTC
TTGGCCATGGTCCCTGGTGTGCAACTCCTCTCTTCAACGCTCTTGGCTGGCTCCT
TGTC AAGTGTCCAAGGTCCCCAGCTCCATAAGTCAGGAAGAAATCCAGGCAAGACG
TGATCTACCAGAACCTGACCCAGCTTAAAGCTGCAGTGGGTGAGCTCTCAGAGAAA
TCCAAGCTGCAGGAGATCTACCAGGAGCTGACCCAGCTGAAGGCTGCAGTGGGTGA
GCTTCCAGAGAAAATCTAAGCAGCAGGAGATCTACCAGGAGCTGACCCGGCTGAAGG
CTGCAGTGGGTGAGCTTCCAGAGAAAATCTAAGATGCAGGAGATCTACCAGGAGCTG
ACTCGGCTGAAGGCTGCAGTGGGTGAGCTTCCAGAGAAAATCTAAGATGCAGGAGAT
CTACCAGGAGCTGACTCGGCTGAAGGCTGCAGTGGGTGAGCTTCCAGAGAAAATCTA
AGCAGCAGGAGATCTACCAGGAGCTGACCCAGCTGAAGGCTGCAGTGGGTGAGCTT
CCAGAGAAAATCTAAGCAGCAGGAGATCTACCAGGAGCTGACCCAGCTGAAGGCTGC
AGTGGGTGAGCTTCCAGAGAAAATCTAAGCAGCAGGAGATCTACCAGGAGCTGACCC
CGGCTGAAGGCTGCAGTGGAAACGCTGTGCCCGCTGCCCGCTGGGAATGGACATT
CTTCCAAAGGAAACTGTACTTCACTGCTAACTCCAGCGGAACCTGGCACGACTCCAT
CACTGCCTGCAAAAGAAAGTGGGGGCCAGCTCGTCGTAATCAAAAGTGTGAGGAGC
AGAACTTCTACAGCTGCAGTCTTCCAGAAAGTAACCGCTTCACTGATGGGACTTT
CAGATCTAAATGAGGAAGGCAATGTGGCAATGGGTGGACGGCTCACCTGTGTGCC
AGCTTCAACCAGTAYTGGAACAGAGGAGAGGCCCAACAACGTTGGGAGGAAGACTG
CGCGGAATTTAGTGGCAATGGCTGGAATGACGACAAATGTAATCTTGCCAAATCTG
GATCTGCAAAAAGTCCGACGCTCCTGCTCCAGGATGAAGAACAAGTTTCTTCTCC
AGCCCCTGCCACCCCAACCCCTCCTGCG (SEQ. ID. NO. 10)

Fig. 12

Gorilla

ATGAGTACTCAAGGAACCAAGACTGCAGCAGCTGGGCCTCCTGGAGGAGGAACA
GCTGAGAGGCCCTTGGATTCCGACAGACTCAGGCTACAAGAGCTTAGCAGGGTGTCTC
TTGGCCATGGTCCCTGGTGTGCAACTCCTCTCCTTACAGCTCTTGGCTGGCTCCT
TGTCCAAAGTGTCCAAAGGTCCCAAGCTCCATAAGTCAGGAACAATCCAGGCAAGACG
CGATCTACCAGAACCTGACCCAGTTTAAAGCTGCAGTGGTGAGCTCTCAGAGAAA
TCCAAAGCTGCAGGAGATCTATCAGGAGTGAACCCAGCTGAAGGCTGCAGTGGGTGA
GCTTCCAGAGAAATCTAAGCAGCAGGAGATCTACCAGGAGCTGAGCCAGCTGAAGG
CTGCAGTGGTGAGCTTCCAGAGAAATCTAAGCAGCAGGAGATCTACCAGGAGAT
ACCCGGCTGAAGGCTGAGTGGTGAGCTTCCAGAGAAATCTAAGCAGCAGGAGAT
CTACCAGGAGTGAACCCGGCTGAAGGCTGCAGTGGTGAGCTTCCAGAGAAATCTA
AGCAGCAGGAGATCTACCAGGAGCTGAGCCAGCTGAAGGCTGCAGTGGTGAGCTT
CCAGAGAAATCTAAGCAGCAGGAGATCTACCAGGAGCTGAGCCAGCTGAAGGCTGC
AGTGGTGAGCTTCCAGAGAAATCTAAGCAGCAGGAGATCTACCAGGAGCTGACC
CAGCTGAAGGCTGCAGTGGAAACGCTGTGCCCGCTGCCCTGGGAATGGACATT
CTTCCAAAGGAACCTGTTACTTCACTGCTAATCTCCAGCGGAATGGCACGACTCCAT
CACCGCTGCCAAAGTGGGGCCAGCTCGTCAATCAAAAGTGTGAGGAGCTT
AGAACTTCTACAGCTGCAGTCTTCCAGAAAGTAAACCGCTTCACTGATGGGACTTT
CAGATCTAAATCATGAAGCACGTGGCAATGGGTGACGGCTCACCTCTGTGCCC
AGCTTCGAGCAGTATTGGAACAGAGGAGAGCCCAACACGTTGGGAGGAAGACTG
CGCGGAATTTAGTGGCAATGGCTGGAAACGATGACAAATGTAATCTTGCCAAATCTG
GATCTGCAAAAAGTCTGACGCTCCTGCTCCAGGATGAAGAACAGTTTCTTCTCC
AGCCTCTGCCACCCCAACCCCCCTCCTGCG (SEQ. ID. NO. 11)

Fig. 13

1	ctccagacct	accagaaaag	atgcccggat	ggatccctgca	gctccgtggc	ttttctggga
61	agcagcggcc	cctgctctca	agagaccctg	gctcctgatg	gtggccccc	ggttgccagc
121	tgggtgctagg	gactcaggac	agtttccag	aaaaggccaa	gcgggcagcc	cctccagggg
181	ccgggtgagg	aagctggggg	gtgcccaggc	cacactgggt	ccctgaaccc	cctgcttggg
241	tacagtgcag	ctcctcaagt	ccacagacgt	gggcccggc	agcctcctgt	acctgaagga
301	aatcggccgt	ggctggttcg	ggaaggtgtt	cctgggggag	gtgaactctg	gcacagcag
361	tgcccagggtg	gtggtgaagg	agctgcaggc	tagtgccagc	gtgcaggagc	agatgagtt
421	cctggaggag	gtgcagccct	acagggccct	gaagcacagc	aacctgctcc	agtgcctggc
481	ccagtgcgcc	gaggtgacgc	cctacctgct	ggtgatggag	ttctgcccac	tgggggacct
541	caagggtac	gtcgggagct	gcccgggtgg	ggagtccatg	gctcccagcc	cccggacct
601	gcagcgcatg	gcctgtgagg	tggcctgtgg	cgtcctgcac	cttcacgcga	acaatttcgt
661	gcacagcgac	ctggccctgc	ggaactgcct	gctcacggct	gacctgacgg	tgaagattgg
721	tgactatggc	ctggctcact	gcaagtacag	agaggactac	ttcgtgactg	ccgaccagct
781	gtgggtgcct	ctgcgctgga	tgcgcccaga	gctggtggac	gaggtgcata	gcaacctgct
841	cgtcgtggac	cagaccaaga	gcgggaatgt	gtggtccctg	ggcgtgacca	tctgggagct
901	ctttgagctg	ggcacgcagc	cctatcccca	gcactcggac	cagcaggtgc	tggcgtacac
961	ggtccgggag	cagcagctca	agctgccc	gccccagctg	cagctgacct	tgtcggacct
1021	ctggtacgag	gtgatgcagt	tctgctggct	gcagcccag	cagcggccca	cagccgagga
1081	ggtgcacctg	ctgctgtcct	acctgtgtgc	caagggcg	accgaagcag	aggaggagtt
1141	tgaacggcgc	tggcgtcttc	tgcggcccgg	cgggggcggc	gtggggcccg	ggcccgggtg
1201	ggcggggccc	atgctggg	gcgtgggtga	gctgcgcgct	gcctcgtcct	tcccgtgct
1261	ggagcagttc	gcgggcgacg	gcttccacgc	ggacggcgac	gacgtgctga	cgggtgaccga
1321	gaccagccga	ggcctcaatt	ttgagtacaa	gtgggaggcg	ggccgcggcg	cggaggccctt
1381	cccggccacg	ctgagccctg	gccgcaccgc	acgcctgcag	gagctgtg	cccccgacgg
1441	cgcgcccccg	ggcgtggttc	cgggtgctcag	cgcgcacagc	ccgtcgttgg	gcagcgagta
1501	cttcacccgc	ctagaggagg	ccgcacccgc	cgcggcccac	gacctgact	gcgcgggctg
1561	cgcgccccag	ccacctgcca	ccgcggacca	ggacgacgac	tctgacggca	gcaccgccc
1621	ctcgtggcc	atggagccgc	tgctgggcca	cgggcccacc	gtcgacgtcc	cctggggccg
1681	cggcgaccac	tacctcgcga	gaagcttggc	gcgggaccgc	ctctgcccct	cacgctctcc
1741	ctgcgccctg	gcggggcccc	tgagtctggc	ggaggaggga	gcggaggatg	cagactgggg
1801	cgtggccg	ttctgtcctg	ccttcttcga	ggacccactg	ggcacgtccc	ctttggggag
1861	ctcagggg	cccccgctgc	cgtgactgg	cgaggatgag	ctagaggagg	tgggagcg
1921	gagggccg	cagcgcgggc	actggcgctc	caacgtgtca	gccaacaaca	acagcggcag
1981	ccgctgtcca	gagtcctggg	accccgtctc	tgcgggctgc	cacgctgagg	gctgcccag
2041	tccaaagcag	accccacggg	cctcccccca	gcccgggtac	cctggagagc	ctctgcttgg
2101	gctccaggca	gcctctgccc	aggagccagg	ctgctgccc	ggcctccctc	atctatgctc
2161	tgcccagggc	ctggcacctg	ctccctgctc	ggttacaccc	tctggacag	agacagccag
2221	tagtgggggt	gaccaccgcg	aggcagagcc	caagcttgcc	acggaggctg	agggcactac
2281	cggaccccg	ctgccccttc	cttccgtccc	ctcccatcc	caggaggagg	ccccacttcc
2341	ctcggaggag	gccagtgc	ccgacgccc	tgatgcctg	cctgactctc	ccacgcctgc
2401	tactggtggc	gaggtgtctg	ccatcaagct	ggcttctg	ctgaatggca	gcagcagctc
2461	tcccagagtg	gaggcaccca	gcagtgagga	tgaggacacg	gctgaggcca	cctcaggcat
2521	cttcaccgac	acgtccagcg	acggcctgca	ggccaggagg	ccggatgtgg	tgccagcctt
2581	ccgctctctg	cagaagcagg	tggggacccc	cgactccctg	gactccctgg	acatcccgctc
2641	ctcagccagt	gatggtggct	atgaggtctt	cagcccgctg	gccactggcc	cctctggagg
2701	gcagccgcga	gcgctggaca	gtggctatga	caccgagaac	tatgagtccc	ctgagtttgt
2761	gctcaaggag	gcgcaggaag	ggtgtgagcc	ccaggccctt	gcggagctgg	cctcagaggg
2821	tgagggcccc	gggcccgaga	cacggctctc	cacctccctc	agtggcctca	acgagaagaa
2881	tccctaccga	gactctgcct	acttctcaga	cctcgaggct	gaggccgagg	ccacctcagg
2941	cccagagaag	aagtgcggcg	gggaccgagc	cccggggcca	gagctggggc	tgccgagcac
3001	tgggcagccg	tctgagcagg	tctgtctcag	gcctgggggt	tccggggagg	cacaaggctc

Figure 14A

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3061 tggccccggg gaggtgctgc cccactgct gcagcttgaa gggtcctccc cagagcccag
3121 cacctgcccc tcgggcctgg toccagagcc tccggagccc caaggcccag ccaagggtgcg
3181 gcctggggccc agccccagct gctcccagtt tttcctgctg accccggttc cgctgagatc
3241 agaaggcaac agctctgagt tccaggggcc ccaggactg ttgtcagggc cggccccaca
3301 aaagcggatg gggggcccag gcacccccag agccccactc cgcctggctc tgcccggcct
3361 ccctgcgggc ttggaggggc ggccggaggga ggaggaggag gacagtgagg acagcgacga
3421 gtctgacgag gagctccgct gctacagcgt ccaggagcct agcgaggaca gcgaagagga
3481 ggcgccggcg gtgcccgtgg ttggtggctga gagccagagc gcgcgcaacc tgcgcagcct
3541 gctcaagatg cccagcctgc tgtccgagac cttctgcgag gacctggaac gcaagaagaa
3601 ggccgtgtcc ttcttcgacg acgtcacogt ctacctcttt gaccaggaaa gccccaccg
3661 ggagctcggg gagcccttcc cgggcgccaa ggaatcgccc cctacgttcc ttagggggag
3721 ccccggtctc cccagcgccc ccaaccggcc gcagcaggct gatggctccc caaatggctc
3781 cacagcgga gaggggtggtg ggttcgctg ggacgacgac ttcccgtga tgacggccaa
3841 ggcagccttc gccatggccc tagaccggc cgcacccgcc ccggtgcgc ccacggccac
3901 gcccgtctcc ttctcgctc tcacgggtgtc gcccgcgcc acgtcccgtc tctccatcac
3961 gcacgtgtct gactcggacg ccgagtccaa gagaggacct gaagctgggt cggggggtga
4021 gagtaaagag gcttgagacc tgggcagctc ctgccccca aggetggcgt caccggagcc
4081 cctgccaggc agcagcgagg atggtgaccg agaaggtggg gaccacgtcc tgggtggctgt
4141 tggcagcaga ttcaggtgcc tctgccccac gcggtgtcct ggagaagccc gtgggatgag
4201 aggccttggg ttgtagatcg gccatgctcc gcccagagg cagaattcgt ctgggctttt
4261 aggttgcgtg ctagcccctg ggggcgcctg gagccacagt ggggtgtctgt acacacatac
4321 aactcaaaa ggggccagt cccctgggca cggcgccccc caccctctgc cctgctgcc
4381 tggcctcgga ggaccgcgt gcccatccg gcagctcctc cgggtgtgctc acaggacact
4441 taaaccagga cgaggcatgg ccccgagaca ctggcaggtt tgtgagcctc tcccacccc
4501 ctgtgcccc acccttgctt ggttcctggt ggctcagggc aaggagtggc cctgggcgcc
4561 cgtgtcggtc ctgtttccgc tgcccttata tcaaagtccg tggctgtttc cccttactg
4621 actcagctag acccgtaagc ccacccttcc cacagggaac aggetgctcc cacctgggtc
4681 ccgctgtggc cacggtgggc agccaaaag atcaggggtg gaggggcttc caggctgtac
4741 tcctgccccg tgggccccgt tctagagggt cccttggcag gaccgtgcag gcagctcccc
4801 tctgtggggc agtatctggt cctgtgcccc agctgccaaa ggagagtggg ggccatgccc
4861 cgcagtcagt gttggggggc tcctgcctac agggagaggg atggtgggga aggggtggag
4921 ctgggggagc ggcagcacag ggaatatatt tgtaactaac taactgctgt gggtggagcg
4981 aatggaagtt gggtgatatt aagtatttgt tgccaaagag atgtaaagtt tattgttgct
5041 tcgcaggggg atttgttttg tgtttgtttt gaggcttaga acgctgggtg aatgttttct
5101 tgttccttgt tttttaagag aaatgaagct aagaaaaaag (SEQ ID NO: 14 and 15)

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Figure 14A (continued)

MQFLEEVQPYRALKHSNLLQCLAQCAEVTPYLLVMEFCPLGDLKGYLRSCRVAESMAP
DPRTLQRMACEVACGVLHLHRNNFVHSDLALRNCLLTADLTVKIGDYGLAHCKYRED
YFVTADQLWVPLRWIAPELVDEVHSNLLVVDQTKSGNVWSLGVTIWELFELGTQYPYQ
HSDQQVLAYTVREQQKLKPKPQLQLTLSDRWYEVMQFCWLQPEQRPTAEEVHLLLSYL
CAKGATEAEEEFERRWRSRPGGGGVGP GPAAGPMLGGVVELAAASSFPLLEQFAGD
GFHADGDDVLTVTETSRGLNFEYKWEAGRGAEAFPATLSPGRTARLQELCAPDGAPPG
VVPVLSAHSPSLGSEYFIRLEEAAPAAGHDPDCAGCAPSPPATADQDDSDGSTAASLA
MEPLLGHGPPVDVPWGRGDHYPRRSLARDPLCPSRSPSPSAGPLSLAEGGAEDADWGV
AAFCPAFFEDPLGTSPLGSSGAPPLPLTGEDELEEVGARAAQRGHWRSNVSANNNSGS
RCPESWDPVSAGCHAEGCPSPKQTPRASPEPGYPGEPLLGLQAASAQEPGCCPGLPHLCS
AQGLAPAPCLVTPSWTETASSGGDHPQAEPKLATEAEGTTGPRLPLPSVPSPSQEGAPLP
SEEASAPDAPDALPDSPTPATGGEVSAIKLASALNGSSSSPEVEAPSEDEDTAEATSGIFT
DTSSDGLQARRPDVVPAFRSLQKQVGTPDSLDSLIPSSASDGGYEVFSPSATGPSGGQP
RALDSGYDTENYESPEFVLKEAQEGCEPQAFaelaseGEGPGPETRLSTSLSGLNEKNPY
RDSAYFSDLEAEAEATSGPEKKCGGDRAPGPELGLPSTGQPSEQVCLRPGVSGEAQGS
PGEVLPPLLQLEGSSPEPSTCPSGLVPEPPEPQGPakVRPGPSPSCSQFFLLTPVPLRSEGN
SSEFQGPPGLLSGPAPQKRMGGPGTPRAPLRLALPGLPAALEGRPEEEEEEDSEDSDSDE
ELRCYSVQEPSSEDEEEAPAVPVVVAESQSARNLRSLLKMPSLLSETFCEDLERKKKAVS
FFDDVTVYLFQESPTRELGEFPGAKESPPTFLRGSPGSPSAPNRPQQADGSPNGSTAEE
GGGFAWDDDFPLMTAKAAAFAMALDPAAPAPAAPTPTPAPFSRFTVSPAPTSRFSITHVS
DSDAESKRGPEAGAGGESKEA (SEQ ID NO:16)

Figure 14B

GCTCCCTGCCTGGTTACACCCTCCTGGACAGAGACAGCCGGTAGTGGGGGTGACCACCCGCAGGCAGAGCC
 CAAGCTTGCCACGGAGGCTGAGGGCACTGCCGGACCCTGTCTGCCCCCTTCCTTCCGTCCCCCTCCCCATCCC
 AGGAGGGAGCCCCACTTCCCTCGGAGGAGGCCAGTGCCCCCTGACGCCCCCTGATGCCCTGCCTGACTCTCCC
 ATGCCTGCTACTGGTGGCGAGGTGTCTGCCATCAAGCTGGCTTCTGTCTGAATGGCAGCAGCAGCTCTCC
 CGAGGTGGAGGCACCCAGCAGCGAGGATGAGGACACGGCTGAGGCCACCTCAGGCATCTTACCCGACACGT
 CCAGCGACGGCCTGCAGGCCGAGAGGCTGGATGTGGTGCCAGCCTTCCGCTCTCTGCAGAAGCAGGTGGGG
 ACCCCCGACTCCCTGGACTCCCTGGACATCCCATCCTCAGCCAGTGATGGTGGCTATGAGGTCTTCAGCCC
 GTCGGCCACTGGCCCCCTCTGGAGGGCAGCCCCGAGCGCTGGACAGTGGCTATGACACCGAGAAGTATGAGT
 CCCCTGAGTTTGTGCTCAAGGAGGCGCAGGAAGGGTGTGAGCCCCAGGCCTTTGAGGAGCTGGCCTCAGAG
 GGTGAGGGCCCCGGCCCCGGGCCCGAGACGCGCTCTCCACCTCCCTCAGTGGCCTCAACGAGAAGAATCC
 CTACCGAGACTCTGCCTACTTCTCAGACCTGGAGGCTGAGGCCGAGGCCGAGGCCACCTCAGGCCACAGAGA
 AGAAGTGCGGCGGGGACCAAGCCCCCGGGCCAGAGCTGGACCTGCCGAGCACTGGGCAGCCGTCTGAGCAG
 GTCTCCCTCAGGCCCTGGGGTTTCCGGGGAGGCACAAGGCTCTGGCCCCGGGGAGGTGCTGCCCCCACTGCT
 GCGGCTTGAAGGATCCTCCCCAGAGCCAGCACCTGCCCTCGGGCCTGGTCCCAGAGCCTCCGGAGCCCC
 AAGGCCAGCCGAGGTGCGGCCTGGGCCAGCCCCAGCTGCTCCAGTTTTTCTGCTGACCCCGGTTCCG
 CTGAGATCAGAAGGCAACAGCTCTGAGTTCCAGGGGCCCCCAGGACTGTTGTGAGGGCCGGCCCCACAAAA
 GCGGATGGGGGGCCTAGGCACCCCCAGAGCCCCACTCCGCTGGCTCTGCCCGGCCCTCCCTGCGGCCTTGG
 AGGGCCGGCCGGAGGAGGAGGAGGAGGACAGTGAGGACAGCGCGAGTCTGACGAGGAGCTCCGCTGCTAC
 AGCGTCCAGGAGCCTAGCGAGGACAGCGAAGAGGAGGCGCCGCGGTGCCCGTGGTGGTGGCTGAGAGCCA
 GAGCGCGCAACCTGCGCAGCCTGCTCAAGATGCCAGCCTGCTGTCCGAGGCCTTCTGCGAGGACCTGG
 AACGCAAGAAGAAGGCCGTGTCTTCTTCGACGACGTACCGTCTACCTCTTTGACCAGGAAAGCCCCACC
 TGGGAGCTCGGGGAGCCCTTCCGGGGCGCCAAGGAATCGCCCCCACGTTCTTAGGGGGAGCCCCGGCTC
 TCCCAGCGCCCCAACCGGCCGACGAGGCTGATGGCTCCCCAAATGGCTCCACAGCGGAAGAGGGTGGTG
 GGTTCGCGTGGGACGACGACTTCCCGCTGATGCCGGCCAAGGCAGCCTTCGCCATGGCCCTAGACCCGGCC
 GCACCCGCCCCGGCTGCGCCACGCCC*****GCTCCCTTCTCGCGCTTCACGGTGTGCCCCGCGCCAC
 GTCCACGTCCCGCTTCTCCATCACGCACGTGTCT (SEQ ID NO:17)

Figure 15A

GCTCCCTGCCTGGTTACACCCTCCTGGACAGAGACAGACGGTAGTGGGGGTGACCACCCGCAGGCAGAGCC
 CAAGCTTGCCACGGAGGCTGAGGGCACTGCCGGACCCCGCTGCCCTTCCTTCCGTCCCCCTCCCCATCCC
 AGGAGGGAGCCCCACTTCCCTCGGAGGAGGCCAGTGCCCCGACGCCCCCTGATGCCCTGCCTGACTCGCCC
 ACGCTGCTACTGGTGGCGAGGTGTCTGCCACCAAGCTGGCTTCCGCCCTGAATGGCAGCAGCAGTCTCC
 CGAGGTGGAGGCACCCAGCAGTGAGGATGAGGACACGGCTGAGGCAACCTCAGGCATCTTACCCGACACGT
 CCAGCGACGGCCTGCAGGCCGAGAGGCAGGATGTGGTGCCAGCCTTCCACTCTCTGCAGAAGCAGGTGGGG
 ACCCCCGACTCCCTGGACTCCCTGGACATCCCGTCCCTCAGCCAGTGATGGTGGCTATGAGGTCTTCAGCCC
 GTCGGCCACGGGCCCTCTGGAGGGCAGCCCCGAGCGCTGGACAGTGGCTATGACACCGAGAAGTATGAGT
 CCCCTGAGTTTGTGCTCAAGGAGGCGCAGGAAGGGTGTGAGCCCCAGGCCTTTGCGGAGCTGGCCTCAGAG
 GCGAGGGC*****CCGGGCCCGAGACGCGGCTCTCCACCTCCCTCAGTGGCCTCAACGAGAAGAATCC
 CTACCGAGATTCTGCCTACTTCTCAGACCTGGAGGCT*****GAGGCCGAGGCCTACCTCAGGCCACAGAGA
 AGAAGTGCGGTGGGGACCAAGCCCCCGGGCCAGAGCTGGGCCTGCCGAGCACTGGGCAGCCGTCTGAGCAG
 GTCTCCCTCAGTCTTGGGGTTTCCGTGGAGGCACAAGGCTCTGGCCCCGGGGAGGTGCTGCCCCCACTGCT
 GCGGCTTGAAGGGTCTCCCCAGAGCCAGCACCTGCCCTCGGGCCTGGTCCCAGAGCCTCCGGAGCCCC
 AAGGCCAGCCGAGGTGCGGCCTGGGCCAGCCCCAGCTGCTCCAGTTTTTCTGCTGACCCCGGTTCCG
 CTGAGATCAGAAGGCAACAGCTCTGAGTTCCAGGGGCCCCCAGGACTGTTGTGAGGGCCGGCCCCACAAAA
 GCGGATGGGGGGCCAGGCACCCCCAGAGCCCCACACCGCCTGGCTCTGCCCGGCCCTCCCTGCGGCCTTGG
 AGGGCCGGCCGGAGGAGGAGGAGGACAGTGAGGACAGCGACGAGTCTGACGAGGAGCTCCGCTGCTAC
 AGCTCCAGGAGCCTAGCGAGGACAGCGAAGAGGAGGCGCCGCGGTGCCCGTGGTGGTGGCTGAGAGCCA
 GAGCGCGCAACCTGCGCAGCCTGCTCAAGATGCCAGCCTGCTGTCCGAGGCCTTCTGCGAGGACCTGG
 AACGCAAGAAGAAGGCCGTGTCTTCTTCGACGACGTACCGTCTACCTCTTTGACCAGGAAAGCCCCACC
 CGGGAGCTCGGGGAGCCCTTCCGGGGCGCCAAGGAATCGCCCCCACGTTCTTAGGGGGAGCCCCGGCTC
 TTCCAGCGCCCCAACCGGCCGACGAGGCTGATGGCTCCCCAAATGGCTCCACAGCGGAAGAGGGTGGTG
 GGTTCGCGTGGGACGACGACTTCCCGCTGATGCCGGCCAAGGCAGCCTTCGCCATGGCCCTAGACCCGGCC
 GCACCCGCCCCGGCTGCGCCACGCCC*****GCTCCCTTCTCGCGCTTCACGGTGTGCCCCGCGCCAC
 GTCC:::CGCTTCTCCATCACGCACGTGTCT (SEQ ID NO:18)

Figure 15B